



## SEQUENCE LISTING

<110> Dixit, et al.  
<120> Interleukin-1 Beta Converting Enzyme Like Apoptotic Protease-6  
<130> PF335D2  
<140> US 09/961,201  
<141> 2001-09-24  
<150> US 09/300,328  
<151> 1999-04-27  
<150> US 08/852,936  
<151> 1997-05-08  
<150> US 60/018,961  
<151> 1996-06-05  
<150> US 60/020,344  
<151> 1996-05-23  
<150> US 60/017,949  
<151> 1996-05-20  
<160> 11  
<170> PatentIn version 3.1  
<210> 1  
<211> 416  
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<213> Homo sapiens  
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Glu Leu Phe Arg Pro His Met Ile Glu Asp Ile Gln Arg Ala Gly Ser  
35 40 45

Gly Ser Arg Arg Asp Gln Ala Arg Gln Leu Ile Ile Asp Leu Glu Thr  
50 55 60

Arg Gly Ser Gln Ala Leu Pro Leu Phe Ile Ser Cys Leu Glu Asp Thr  
65 70 75 80

Gly Gln Asp Met Leu Ala Ser Phe Leu Arg Thr Asn Arg Gln Ala Gly  
85 90 95

Lys Leu Ser Lys Pro Thr Leu Glu Asn Leu Thr Pro Val Val Leu Arg  
100 105 110

Pro Glu Ile Arg Lys Pro Glu Val Leu Arg Pro Glu Thr Pro Arg Pro  
115 120 125

Val Asp Ile Gly Ser Gly Gly Phe Gly Asp Val Gly Ala Leu Glu Ser  
 130 135 140

Leu Arg Gly Asn Ala Asp Leu Ala Tyr Ile Leu Ser Met Glu Pro Cys  
 145 150 155 160

Gly His Cys Leu Ile Ile Asn Asn Val Asn Phe Cys Arg Glu Ser Gly  
 165 170 175

Leu Arg Thr Arg Thr Gly Ser Asn Ile Asp Cys Glu Lys Leu Arg Arg  
 180 185 190

Arg Phe Ser Ser Leu His Phe Met Val Glu Val Lys Gly Asp Leu Thr  
 195 200 205

Ala Lys Lys Met Val Leu Ala Leu Leu Glu Leu Ala Arg Gln Asp His  
 210 215 220

Gly Ala Leu Asp Cys Cys Val Val Val Ile Leu Ser His Gly Cys Gln  
 225 230 235 240

Ala Ser His Leu Gln Phe Pro Gly Ala Val Tyr Gly Thr Asp Gly Cys  
 245 250 255

Pro Val Ser Val Glu Lys Ile Val Asn Ile Phe Asn Gly Thr Ser Cys  
 260 265 270

Pro Ser Leu Gly Gly Lys Pro Lys Leu Phe Phe Ile Gln Ala Cys Gly  
 275 280 285

Gly Glu Gln Lys Asp His Gly Phe Glu Val Ala Ser Thr Ser Pro Glu  
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Asp Glu Ser Pro Gly Ser Asn Pro Glu Pro Asp Ala Thr Pro Phe Gln  
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Glu Gly Leu Arg Thr Phe Asp Gln Leu Asp Ala Ile Ser Ser Leu Pro  
 325 330 335

Thr Pro Ser Asp Ile Phe Val Ser Tyr Ser Thr Phe Pro Gly Phe Val  
 340 345 350

Ser Trp Arg Asp Pro Lys Ser Gly Ser Trp Tyr Val Glu Thr Leu Asp  
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Asp Ile Phe Glu Gln Trp Ala His Ser Glu Asp Leu Gln Ser Leu Leu  
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Leu Arg Val Ala Asn Ala Val Ser Val Lys Gly Ile Tyr Lys Gln Met  
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 ccaggggctg tctacggcac agatggatgc cctgtgtcgg tcgaaaagat tgtgaacatc 180  
 ttcaatggga ccagctgcc cagcctggga gggaagccca agctcttttt catccaggcc 240  
 tgtggtgggg agcagaaaga ccatgggttt gaggtggcct ccacttcccc tgaagacgag 300  
 tccccctggca gtaaccccgga gccagatgcc accccgttcc aggaaggttt gaggaccttc 360  
 gaccagctgg acgccatata tagtttgccc acaccagtg acatctttgt gtctactct 420  
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Met Val Leu Ala Leu Leu Glu Leu Ala Arg Gln Asp His Gly Ala Leu  
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Asp Cys Cys Val Val Val Ile Leu Ser His Gly Cys Gln Ala Ser His  
 20 25 30

Leu Gln Phe Pro Gly Ala Val Tyr Gly Thr Asp Gly Cys Pro Val Ser  
 35 40 45

Val Glu Lys Ile Val Asn Ile Phe Asn Gly Thr Ser Cys Pro Ser Leu  
 50 55 60

Gly Gly Lys Pro Lys Leu Phe Phe Ile Gln Ala Cys Gly Gly Glu Gln  
 65 70 75 80

Lys Asp His Gly Phe Glu Val Ala Ser Thr Ser Pro Glu Asp Glu Ser  
 85 90 95

Pro Gly Ser Asn Pro Glu Pro Asp Ala Thr Pro Phe Gln Glu Gly Leu  
 100 105 110

Arg Thr Phe Asp Gln Leu Asp Ala Ile Ser Ser Leu Pro Thr Pro Ser  
115 120 125

Asp Ile Phe Val Ser Tyr Ser Thr Phe Pro Gly Phe Val Ser Trp Arg  
130 135 140

Asp Pro Lys Ser Gly Ser Trp Tyr Val Glu Thr Leu Asp Asp Ile Phe  
145 150 155 160

Glu Gln Trp Ala His Ser Glu Asp Leu Gln Ser Leu Leu Leu Arg Val  
165 170 175

Ala Asn Ala Val Ser Val Lys Gly Ile Tyr Lys Gln Met Pro Gly Cys  
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Phe Asn Phe Leu Arg Lys Lys Leu Phe Phe Met  
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